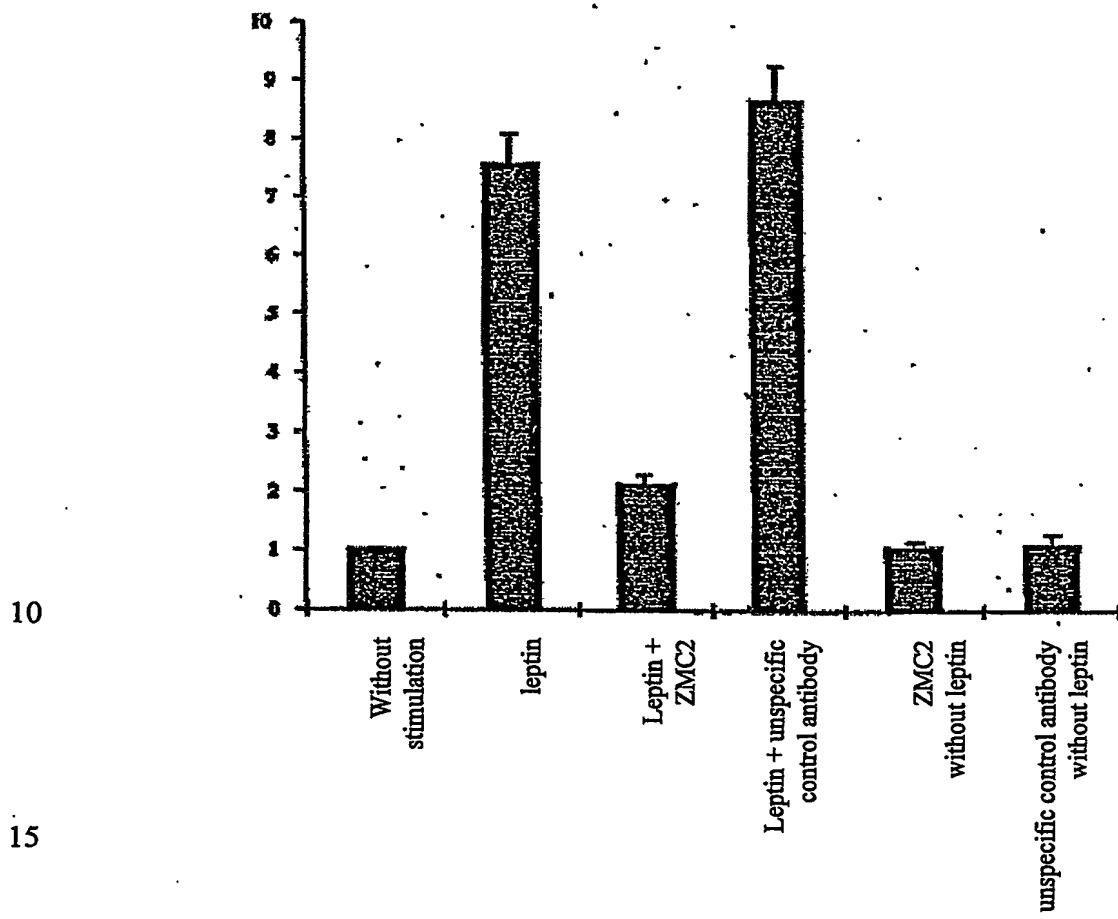


1/20

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Figure 1

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2/20

A)

5 XHNPIPMPPAAAGLLLLAAQPAMAEVMTQSPKFMSTSIGDRVNITCKAT
 NVVRTAVTWYQQKPGQSPQALIFLASNRRHTGVPARFTGSGSGTDFTLTIN
 NVKSEDLADYFCLQHWNYPPLTFGSGTKLEIKRADAAPTVSIFPPSSEQLT
 SGGASVVCFLNNFYPKDINVWKWIDGSERQNGVILNSWTDQDSKDSTYSMS
 STLTLTKDEYERHNSYTCEATHKTSTPIVKSFNRGEC**SRVKRXQSXG
 GPGTPIRPIGXPYYNLSGGGFQ

B)

10 DNA: NANGTCTATAATCCAATACCTATGCCCTACGGCAGCCGCTGGATTGTTATTAC
 +3: X H N P I P M P T A A A G L L L L
 pComb3 vector SacI V_L(x) primer

15 DNA: TCGCTGCCCAACCAGCCATGGCCGAGCTCGTGATGATGACCCAGCTCCAAAAT
 +3: A A Q P A M A E L V M T Q S P K F

20 DNA: TCATGTCACATCAATAGGAGACAGGGTCAATATCACCTGCAAGGCCACTC
 +3: M S T S I G D R V N I T C K A T Q

25 DNA: AGAATGTTCGTACTGCTGTTACCTGGTATCAACAGAAACCAGGGCAGTC
 +3: N V R T A V T W Y Q Q K P G Q S P

30 DNA: CTCAAAGCACTGATTTCTTGGCATCCAACCAGGACACTGGTGTCCCTGCTC
 +3: Q A L I F L A S N R H T G V P A R

35 DNA: GATTACAGGCAGTGGATCTGGGACAGATTTCACTCTCACCAATTAAACAATG
 +3: F T G S G S G T D F T L T I N N V

40 DNA: TGAAATCTGAAGACCTGGCAGATTATTCCTGCTACAAACATTGAAATTATC
 +3: K S E D L A D Y F C L Q H W N Y P

45 DNA: CTCTCACGTTCGGCTGGGGACAAAGTTGGAATAAAACGGGCTGATGCTG
 +3: L T F G S G T K L E I K R A D A A

50 DNA: CACCAACTGTATCCATCTTCCCACCATCCAGTGAGCAGTTAACATCTGGAG
 +3: P T V S I F P P S S E Q L T S G G

55 DNA: GTGCCTCAGTCGTGCTTCTTGAAACAACCTTACCCCAAAGACATCAATG
 +3: A S V V C F L N N F Y P K D I N V

60 DNA: TCAAGTGGAAAGATTGATGGCAGTGAACGACAAATGGCGTCTGAACAGTT
 +3: K W K I D G S E R Q N G V L N S W
 BclI

55 DNA: GGACTGATCAGGACAGCAAAGACAGCACCTACAGCATGAGCAGCACCTCA
 +3: T D Q D S K D S T Y S M S S T L T

55 DNA: CGTTGACCAAGGACGAGTATGAACGACATAACAGCTATACTGTGAGGCCA
 +3: L T K D E Y E R H N S Y T C E A T
 C_L(x) primer

55 DNA: CTCACAAGACATCAACTCACCCATTGTCAAGAGCTTCAACAGGGGAGAGT
 +3: H K T S T S P I V K S F N R G E C
 Stop XbaI NotI KpnI

55 DNA: GTTAGTAAATCTAGAGTTAACGGCCGCAATCGAGGGGGCCGGTACCC
 +3: * * S R V K R P Q S R G G F P V P Q

55 DNA: AATTGCCCCATAGGGGNCCGTATTACAATTCACTGGCGGCCTTTCA
 +3: F A L * G X R I T I H W A A V F X

60 DNA: AN
 +3:

Figure 2

3/20

A)

5 LAXRGGGRKIXFXRETVIMKYLXAYGPAAGLLLLAAQPAMAQVKLLESGP
 GLVAPSESLSITCTISGFSLDDGVSWIRQPPGKLEWLGVIWGGGSTYF
 NSLFKSRLSITRDNSKSQVFLEMDSLQTDAMYYCAKHDGHEMDYWGQ
 GTSVTVSSSCTTPPSVYPLAPGSAQTNMVTLGCLVKGYFPEPVTVWN
 SGSSLSSGVHTFPAVLQSDLYTLSSSVPSSTWPSETVTCNVAPASSTK
 VDKKIVPRDCTSHHHHH*ASLVVAVALHSFVXIKANRRPAX

B)

10 DNA: TTGGCCNCCTCCGGTGGCGGCCGCAAATTNTATTNCAAGGGAGACAGTC
 -1: L A X R G G G R K I X F X R E T V

15 DNA: ATAATGAAATACCTTTNGCCTACGGCCAGCCGCTGGATTGTTATTACTC
 -1: I M K Y L X A Y G P A A G L L L L
 pComb3 vector XbaI V_b primer
 DNA: GCTGCCAACAGCCATGGCCCAGGTGAAACTGCTCGAGTCAGGACCTGGC
 -1: A A Q P A M A Q V K L L E S G P G

20 DNA: CTGGTGGCCCTCAGAGAGCCTGTCCATCACATGCACTATCTCAGGGTTC
 -1: L V A P S E S L S I T C T I S G F

25 DNA: TCATTAACCGACGATGGTGTAAAGCTGGATTCTGGCAGCCTCCAGGAAAGGGT
 -1: S L T D D G V S W I R Q P P G K G.

30 DNA: CTGGAGTGGCTGGGAGTAATATGGGTGGTGGAAAGCACATACTTAAATTCA
 -1: L E W L G V I W G G G S T Y F N S

35 DNA: CTTTTCAATCCAGACTGAGCATCACCAAGGGACAACCTCTAAAGAGCCAAGTT
 -1: L F K S R L S I T R D N S K S Q V

40 DNA: TTCTTAGAAATGGACAGTCTACAAACTGATGACACAGCCATGTACTACTGC
 -1: F L E M D S L Q T D D T A M Y Y C

45 DNA: GCCAAACATGACGGACACGAGACTATGGACTATTGGGTCAAGGAACCTCA
 -1: A K H D G H E T M D Y W G Q G T S

50 DNA: GTCACCGCTCCTCATCCAAAAGCACACCCCCATCTGTCTATCCACTGGCC
 -1: V T V S S S K T T P P S V Y P L A

55 DNA: CCTGGATCTGCTGCCAAACTAACCTCCATGGTACCCCTGGGATGCCTGGTC
 -1: P G S A A Q T N S M V T L G C L V

60 DNA: AAGGGCTATTCCTGAGCCAGTGACAGTGACCTGGAACTCTGGATCCCTG
 -1: K G Y F P E P V T V T W N S G S L

65 DNA: TCCAGCGGTGTGCACACCTTCCAGCTGTCTGCAGTCTGACCTCTACACT
 -1: S S G V H T F P A V L Q S D L Y T

70 DNA: CTGAGCAGCTCAGTGAATGTCCTCCAGCACCTGGCCAGCGAGACCGTC
 -1: L S S S V T V P S S T W P S E T V

75 DNA: ACCTGCAACGTTGCCACCCGGCCAGCAGCACCAAGGTGGACAAGAAATT
 -1: T C N V A H P A S S T K V D K K I
 C_{h1}(γ1)Primer SpeI His tag Stop
 DNA: GTGCCAACGGATTGTACTAGTCATCATCATCATCATTAAGCTAGCCTA
 -1: V P R D C T S H H H H H H * A S L

80 DNA: GTGGTGGCGGTGGCTCTCCATTCTGTTGTGANGATAAAGGCCAATCGNAGA
 -1: V V A V A L H S F V X I K A N R R

85 DNA: CCTGCNCNA
 -1: P A X

Figure 3

4/20

A)

ATNCTTNTTGTCTTCTATGCGGCCAGCCGGCCATGGCCCAGGTCCAGCTG
CAGGAGTCAGGA~~ACTGAAGTGGTAAAGCCTGGGCTTCAGTGAAGTTGTCCT~~
GCAAGGCTTCTGGCTACATCTTACAAGTTATGATATA~~AGACTGGGTGAGGCAG~~
5 ACGCCTGA~~ACAGGGACTT~~GA~~GTGGATTGGATGGATTT~~CCTGGAGAGGGGA
GTACTGAATACAATGAGAAGTTCA~~AAGGGCAGGGCACACTGAGTGTAGACAA~~
GTCCTCCA~~GCACAGCCTATATGGAGCTCACTAGGCTGACATCTGAGGACTCTG~~
CTGTCTATTCTGTGCTAGAGGGACTACTATAGGGCTACTTTGACTTGTGGG
10 ~~GCCAAGGGACCACGGTCACCGTCTCCTCATGTGGAGGCAGGTTCAAGGCAGG~~
TGGCTCTGGCGGTGGCGGACTGACATTGAGCTACCCAGTCTCCAGCAATCA
TGTCTGCATCTCCAGGGGAGAGGGTCACCATGACCTGCAGTGCCAGCTC
AAGTATACGTTACATATATTGGTACCAACAGAACGCTGGATCCTCCCCCA
15 GACTCCTGATTATGACACATCCAACGTGGCTCCTGGAGTCCCTTTCGC
TTCAGTGGCAGTGGGTCTGGGACCTCTTATTCTCTCACAA~~CTCAACACCAGAAT~~
GGAGGCTGAGGATGCTGCCACTTATTACTGCCAGGAGTGGAGTGGTTAT
CCTCTCACGTTGGCTCGGGCACCAAGCGGGAAATCAAACGGCGGCCGC
20 AGGTGCGCCGGTGCCGTATCCGGATCCGCTGGAACCGCGTGCCGATAGACT-
GTTGAA

20

B)

MAQVQLQES GTEVVKPGASV~~KLSCKASGYIFTSYIDWVRQTPEQGLEWIG~~
WI~~FPGEGST EYNEKFGRATLSVDKSSSTAYMELTRLTSEDSAVYFCARG~~
25 DYYRRYFDLWGQGTTVTVSSGGGGSGGGGGSDIELTQSPA~~IMSASP~~
GERVTMTCSASSIRIYIY~~WYQQKPGSSPRLLIYDTSNVAPGVFRSGSG~~
SGTSYSLTI~~NRMEAEDAATYYCQEWSGYPLTFGSGTKREIKRAAAGAPVP~~
YPDPLEPR

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35

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Figure 4

5/20

A)

5 tgcgtccccaccagccATGgcccaggtaaaactgctcgagtcaaggacctggcctgg
 cgcctcagagagcctgtccatcacatgcactatctcagggtctcattaaccgacatg
 gtgtaaqctggattcggcagcctccaggaaagggtctggagtggctggagtaatatgg
 qtggtgaaagcacatacttaattcactttcaaatccagactgagcatcaccaggaca
 actctaagagccaagtttcttagaaatggacagtctacaaactqatgacacacagccatgt
 actactqgcggccaaacatqacggacacqagactatggactattgggtcaaggAACCTCAG
 10 tcaccgtctcctcatccaaaacqacacccccatctgtctatccactggccctggatctg
 ctgccccaaactaactccatggtggactctggatccctgtccagcgggtgtgcacacccatgg
 cagtgaagtggactctggatccctgtccagcgggtgtgcacacccatgg
 tcctgcagtctgacctctacactctgagcagctcagtgtactgtccctccagcacctgg
 15 ccagcgagaccgtcacctgcaacgttqcccacccggccagcagcacaagggtggacaaga
 aaattgtgcccaggattgtacttagtggtggcggaggtagtggtggcggaggtagcgg
 qcgagggtctqgtggcggagggtccgaaattctcgaggtggccatccaaaagtccaaag
 atgacacccaaaccctcatcaagacaattgtcaccaggatcaatgacattcacacacgc
 agtcagtctccctccaaacagaaagtccgggtttggacttcatctggctccacccca
 20 tcctgaccttattcaagatggaccagacactggcagtctaccaacagatcctcaccagta
 tgccttccagaaacgtgatccaaatatccaaacgacactggcaggatccctccggatcttc
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 acagcctgggggtgtcctggaaagcttcaggctactccacagagggtggtggccctgagca
 ggctgcaggggctctgtcaggacatgctgtggcagctggacactcagccctgggtgcacta
 25 gtcatcatcatcatcatcatTAActagcttagtggtggcgggtggctctcca

B)

30 Maqvkllesgpqlvapseslsitctisqfslddgvswirappgkglewlgiwgggsty
 fnslfkssltsitrdnsksqvflemdslgtddtamyycahdghetmdywgggtsvtvsss
 kttppsvyplapqsaqtnsmvtlgclvkgyfpepvtnsgslssgvhtfpavlqsd
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 gqseflevpigkvqddtktliktrindishtqsvsskqkvtgldfipglhpiltls
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 35 leasgystevvalsrlqgslqdmilwqldlspgctshhhhh

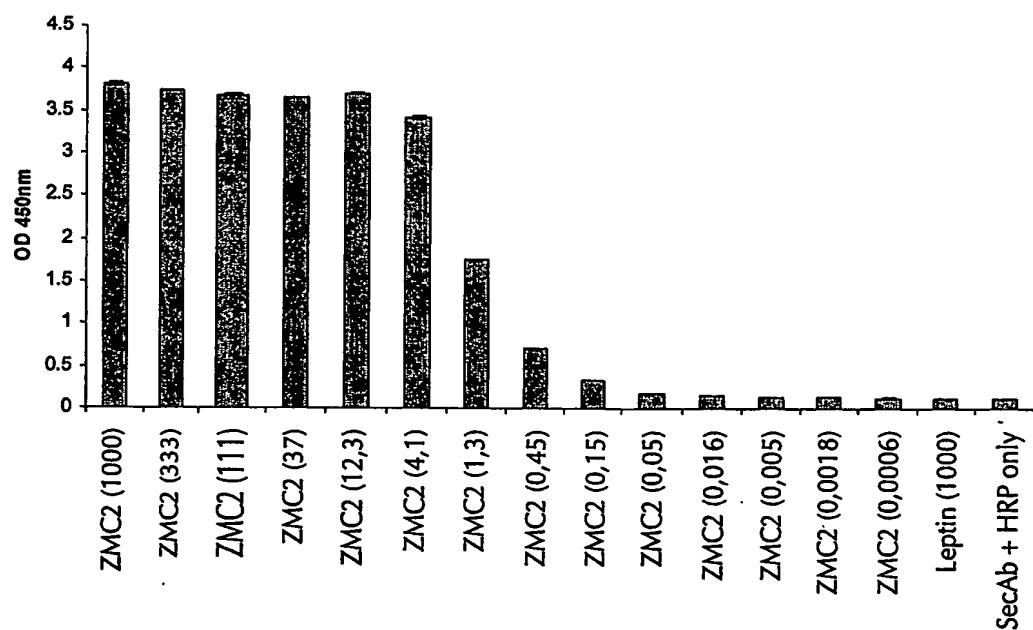
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Figure 5

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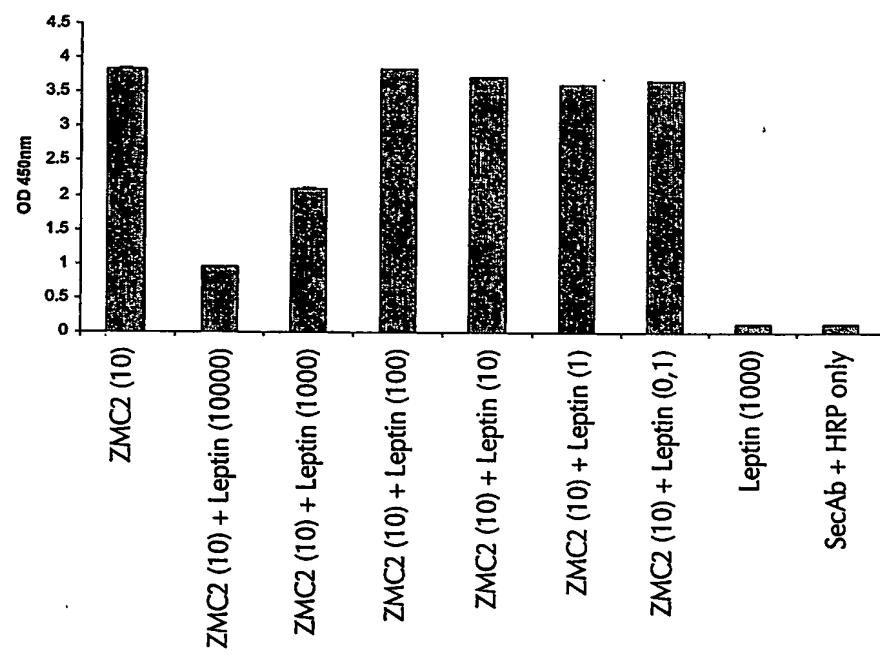
Figure 6

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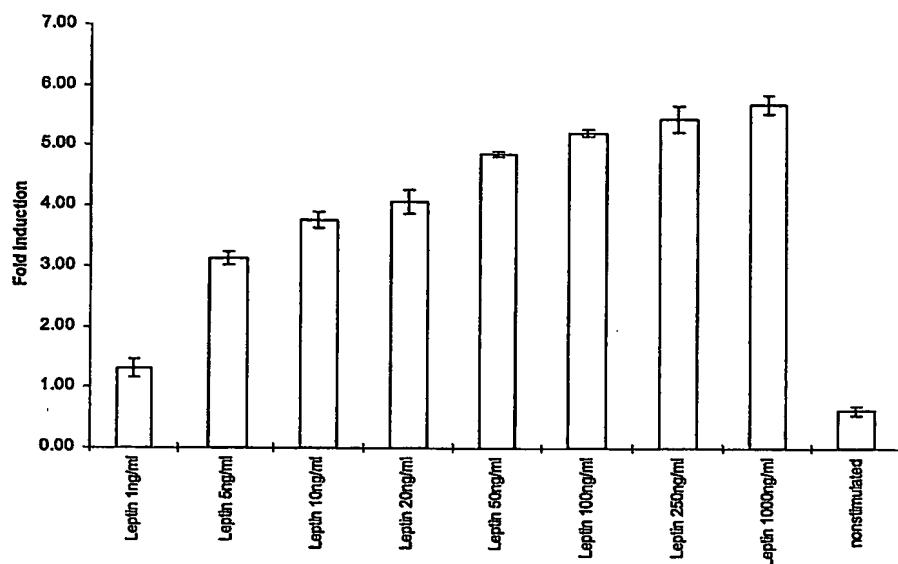
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Figure 7

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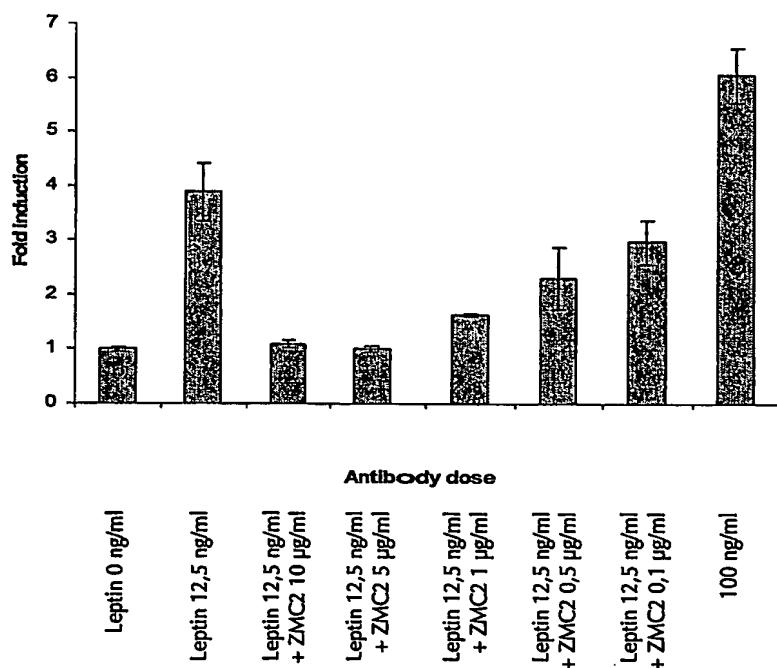
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Figure 8

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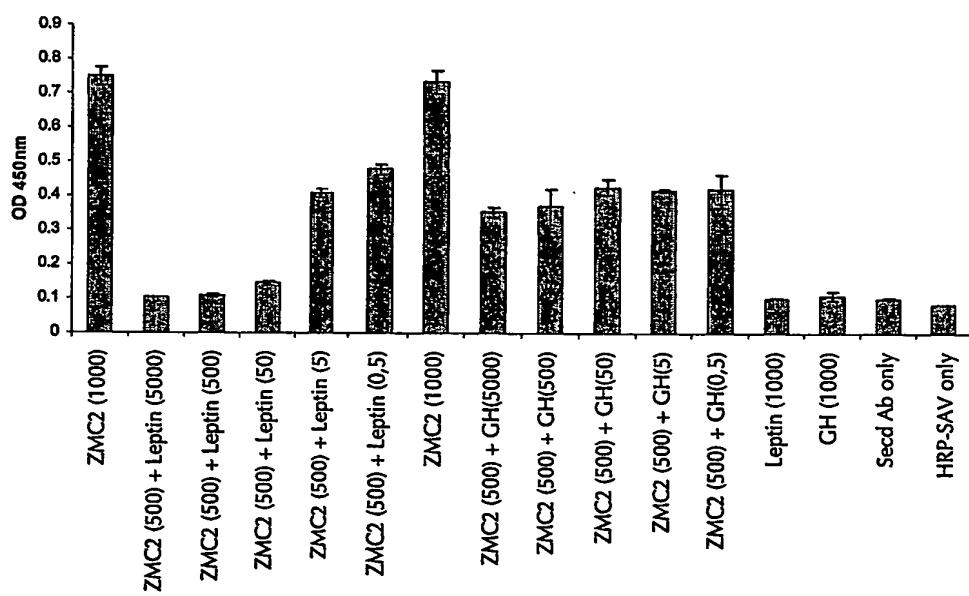
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Figure 9

10/20

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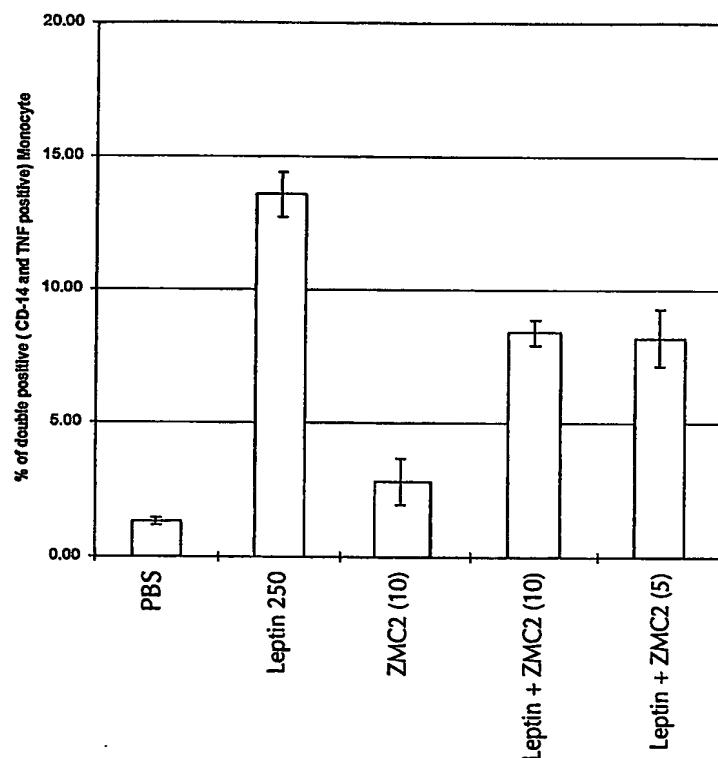
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Figure 10

11/20

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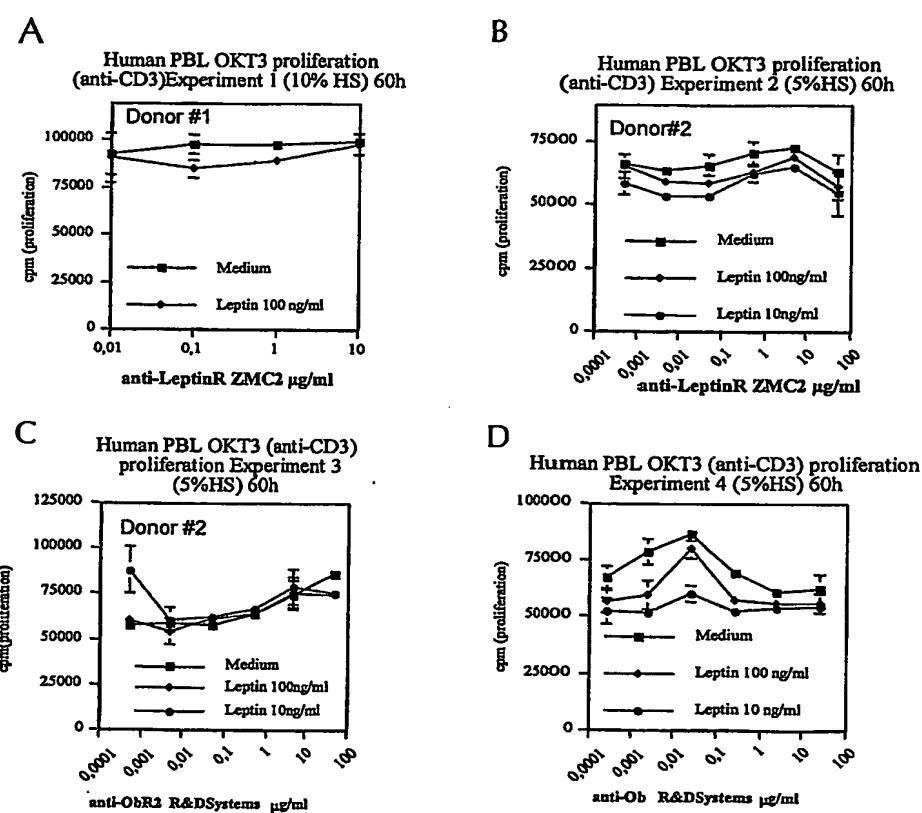
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Figure 11

12/20

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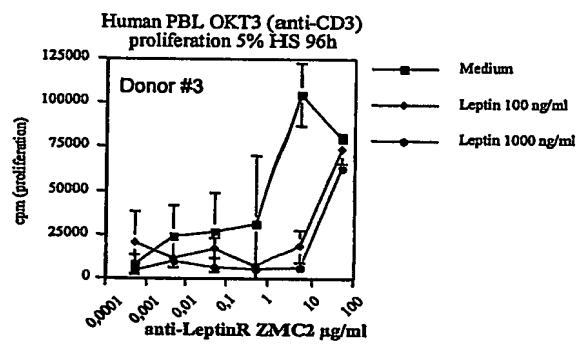
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Figure 12

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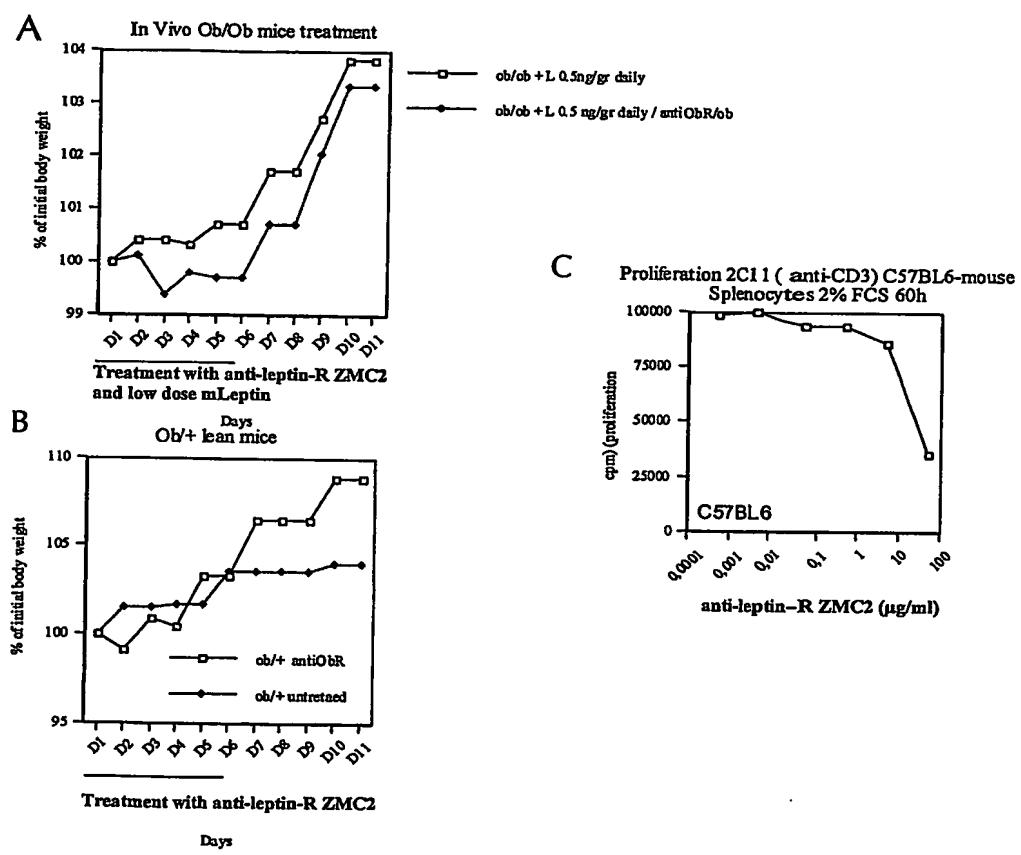
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Figure 13

14/20

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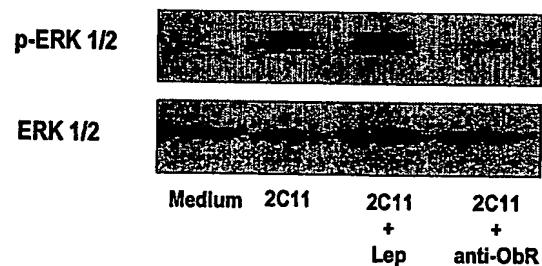
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Figure 14

15/20

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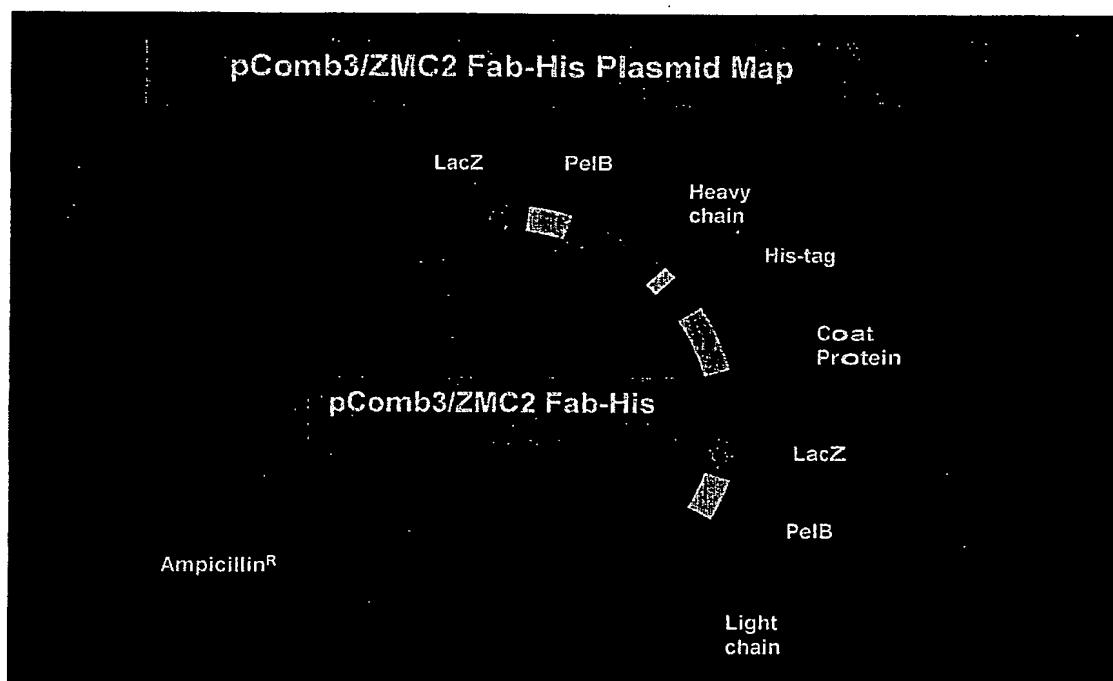
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Figure 15

16/20

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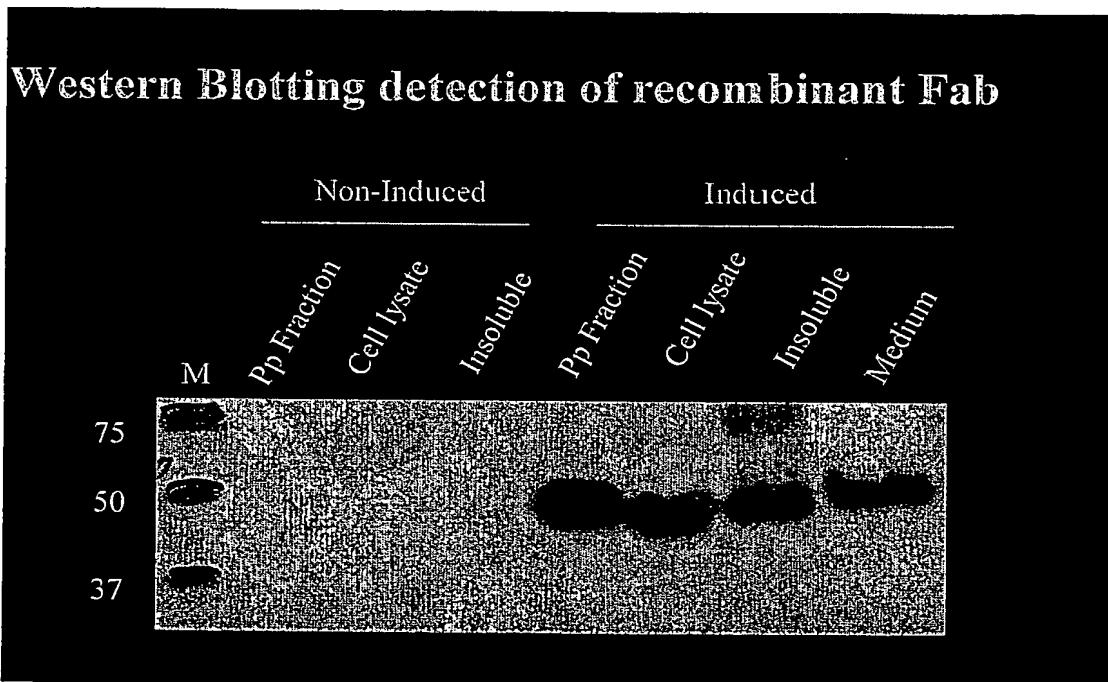
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Figure 16

17/20

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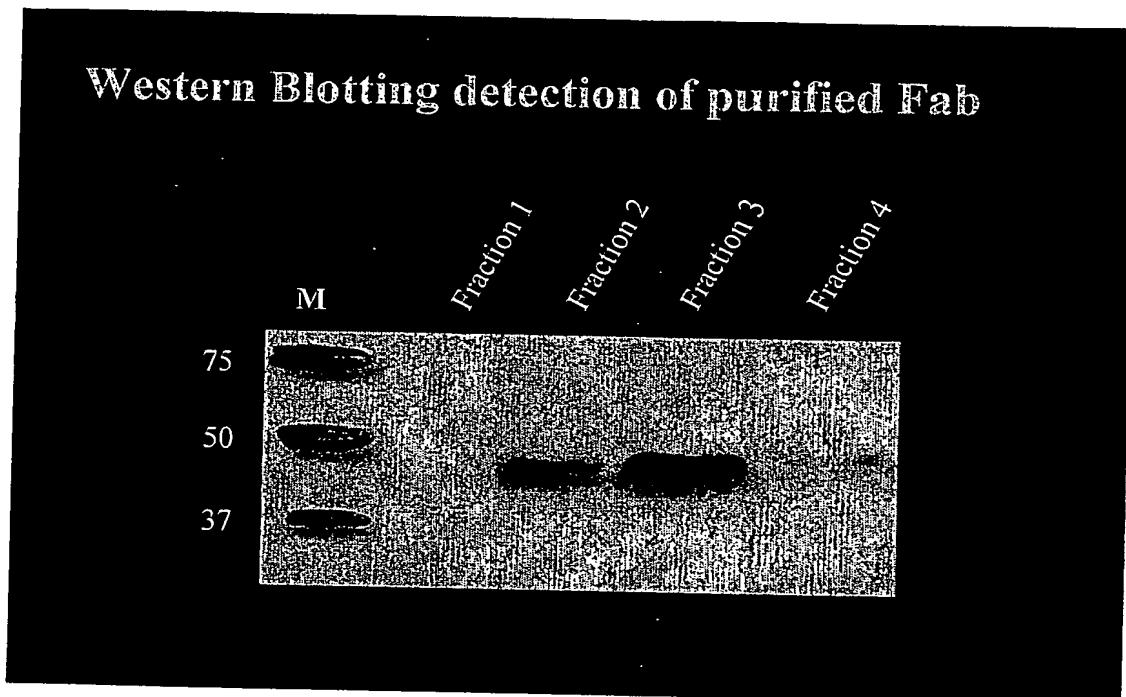
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Figure 17

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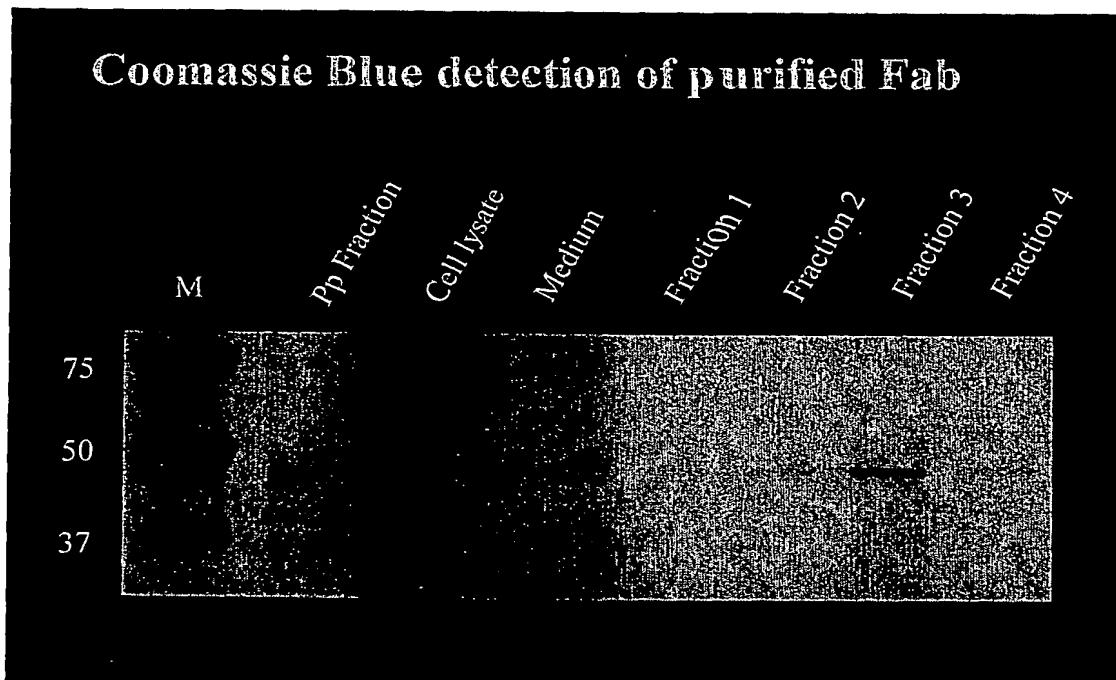
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Figure 18

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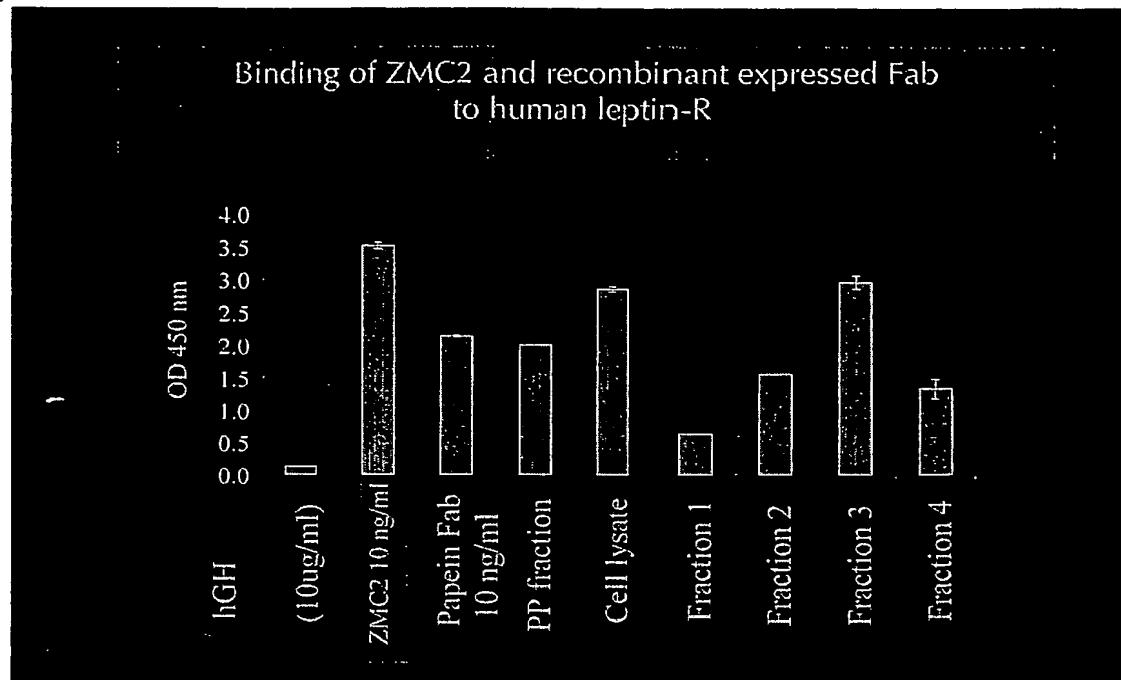
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Figure 19

20/20

A)



B)

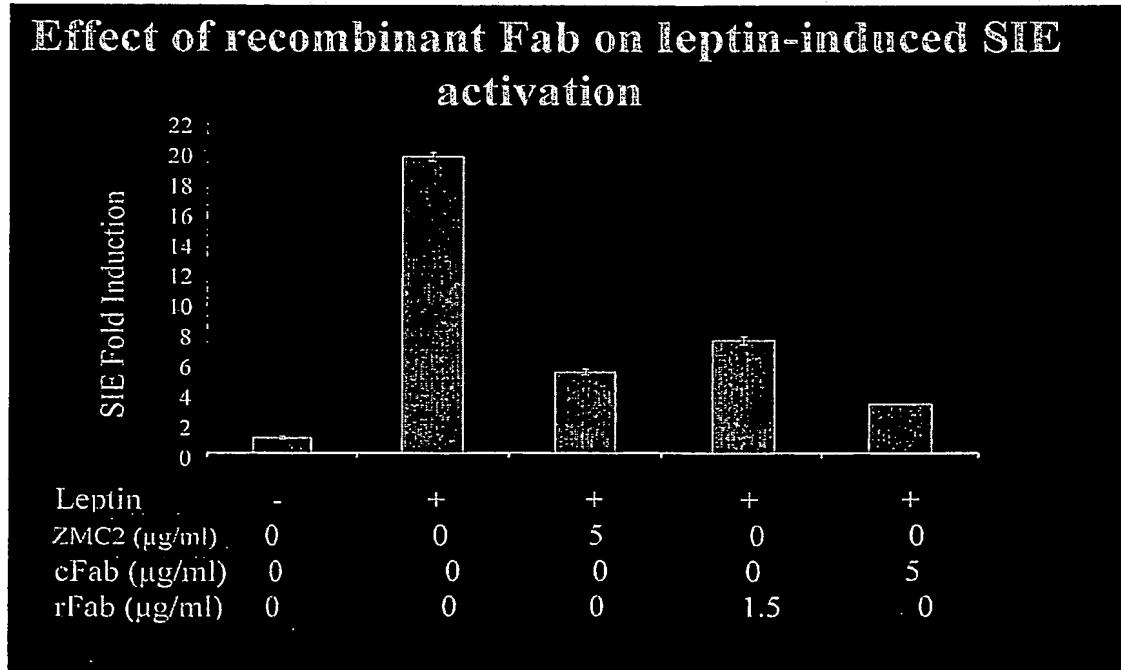


Figure 20